

1 Chitosan conjugated cyclodextrin nanocomposite loaded with antibiotic-adjuvant
2 combinations remediates multi-drug resistant *Staphylococcus aureus* infection in CD-1 mice
3 model of bovine mastitis

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Supplementary information

Supplementary information 1: Assessment of CF, CPZ, and TA for antibacterial synergism

A three-dimensional checkerboard assay, as previously described by Stein *et al.*, was conducted to assess the antibacterial efficiency of the combination (CF, TA, and CPZ) ¹. CF, CPZ, and TA concentrations were selected based on the pre-determined MICs against Sa1158c. Tenfold, six-fold, and seven-fold serial dilutions of CF, CPZ, and TA, respectively, were performed in a 96-well plate. Ten μL of Sa1158c culture maintained at 0.5 McFarland standard (1.5×10^8 cells/ mL) in MHB broth was added further. The plates were incubated at 37 °C for 18 h, and 30 μL of resazurin solution (0.5% of 100X resazurin in PBS buffer (pH-7.4)) was added to each well, and the plates were incubated for 2 h under mild shaking. The fluorescent intensity (530/590 nm) was measured using a plate reader (SpectraMax-i3X, Molecular Devices, USA).

The fractional inhibitory concentration index (FICI) was determined using the following formula:

$$\text{FICI}_{\text{CF/TA/CPZ}} = (\text{MIC}_{\text{CF(combination)}}/\text{MIC}_{\text{CF(alone)}}) + (\text{MIC}_{\text{CPZ(combination)}}/\text{MIC}_{\text{CPZ(alone)}}) + (\text{MIC}_{\text{TA(combination)}}/\text{MIC}_{\text{TA(alone)}})$$

FICI < 0.8, 0.8 < FICI < 4, and FICI \geq 4 were considered synergism, additive or indifference, and antagonism effects, respectively.

Supplementary information 2: Physicochemical characterization of the particles

An attenuated total reflectance-Fourier transform infrared (ATR-FT-IR) (ALPHA-P, Bruker, Billerica, MA, USA) was used to assess the surface functional groups of the particles ². Five microliters of the particle suspension were dropped on the ATR probe and allowed to dry. A wavelength range of 400–4000 cm^{-1} , with a resolution of 4 cm^{-1} , and 32 scans were used to obtain the FT-IR spectrum. Qualitative analysis was performed using the OMNIC 8.2.0.387 software.

45 The FTIR spectra of CPZ-CD-TA and CH Np-CF are provided in **Figure 1a**. The absorbance
46 at 1151 cm^{-1} in CH Np-CF is characteristic of the asymmetric vibration of the b-(1-4) glycosidic
47 bond of chitosan. The low-intensity peaks at 2935 and 2887 cm^{-1} are attributed to stretching
48 vibrations of methylene groups in the polymeric chain. The peaks at 1072 cm^{-1} and 1531 cm^{-1}
49 represent C-O-C- stretching vibrations and -NH_2 bending vibration peaks, respectively. The peak
50 at 1030 cm^{-1} shows the characteristic of P=O stretching vibration from phosphate groups, while the
51 peak at 1628 cm^{-1} is attributed to the electrostatic cross-linking between the phosphate group of TPP
52 and the ammonium group of chitosan, suggesting the formation of CH Np ^{3,4}. CF in CH Np-CF was
53 evident from the additional signals in 1763 , 1635 , 1384 , and 1287 cm^{-1} corresponding to C=O
54 stretching, -NH_2 bending, C-N stretching, and C-O stretching, respectively. CD Np in CPZ-CD-TA
55 was evident from the characteristic bands of saccharides: 2932 cm^{-1} (C-H stretching vibration),
56 1651 cm^{-1} (O-H bending vibration), and 1154 cm^{-1} (C-O vibration) ⁵. The signal at 857 cm^{-1}
57 corresponded to the α -type glycosidic bond, characteristic of CD Np formed by glucopyranose units
58 through the α -1,4-glycosidic bond ⁵. The peak at 2973 cm^{-1} corresponds to the anti-symmetric
59 vibration of a methyl group, indicating the existence of the hydroxypropyl group in CD Np ⁵. CPZ
60 in CPZ-CD-TA was evident from the signals 1606 and 1651 cm^{-1} (phenyl rings), 1510 cm^{-1} (C-H
61 deformation), and 752 cm^{-1} (aromatic C-H bending) ⁶. TA showed an adsorption band at
62 approximately $3000\text{--}3500\text{ cm}^{-1}$ belonging to the stretching vibration of O-H groups. The absorption
63 band at 1187 cm^{-1} belongs to the bending vibration of O-H groups. The skeleton vibration band of
64 benzene rings appeared at $1443\text{--}1704\text{ cm}^{-1}$ ⁷.

65

66 A Scanning Electron Microscope (SEM) (FEG Quanta 450 Field Emission Scanning
67 Electron Microscope, USA) was used to analyse the surface morphology of the particles ⁸.

68 Suspensions of particles (50 µg/mL) were dropped onto aluminum mounts, dried at room
69 temperature, and coated with platinum before acquiring SEM images. ImageJ software was used to
70 measure the size of particles. The hydrodynamic size and surface charge of the particles (50 µg/mL)
71 were measured by Dynamic Light Scattering (DLS) (NanoBrook Omni instrument, Brookhaven,
72 USA) at 25 °C ⁷. The particles were loaded into a pre-rinsed folded capillary cell. A 100 V was
73 applied to measure the zeta potential.

74 CD Np had a typical nano-sized spherical morphology ⁹, while CPZ-CD-TA showed
75 aggregation of particles forming a chain-like arrangement (**Figures 1b-g**). The size of CPZ-CD-TA
76 ranged between ~120-260 nm. DLS studies indicated the hydrodynamic size of CD Np and CPZ-
77 CD-TA to be ~135 nm and a zeta potential of ~-11 mV and ~-9.50 mV, respectively (**Table 1**). The
78 weaker zeta potential might have led to the aggregation of these nanoparticles due to the Van Der
79 Waals attractive forces ⁷. CH Np was uniformly distributed and had a typical spherical structure
80 with a size range between ~114-159 nm. Similarly, CH Np-CF showed a spherical morphology and
81 was of variable sizes (~92-229 nm).

82

83 **Supplementary information 3: *In vitro* antibacterial efficiency of particles**

84 The *in vitro* antibacterial efficiency of the particles was determined using the broth
85 microdilution method ¹⁰. Briefly, the particles (250 µg/mL) were subjected to ten-twofold serial
86 dilution in 100 µL of MHB media in a sterile 96-well plate. Bacterial culture of 10 µL taken from
87 cell suspension with OD adjusted to 0.5 McFarland standard (1.5×10^8 cells/mL) was added to the
88 wells. The plate was incubated at 37 °C for 18 h. The colony-forming units (CFU) were enumerated
89 using a drop plate culturing method, as detailed earlier ^{10, 11}. For this, 10 µL from the treated wells
90 were subjected to seven-fold serial dilutions in 100 µL of PBS (1X, pH 7.4) in a new 96-well plate.

91 Subsequently, 10 μ L of the cell suspension (from 10^1 , 10^3 , 10^5 , and 10^7 dilutions) was dropped onto
92 one end of the rectangular TSA plates and allowed to flow down vertically. These plates were further
93 incubated at 37 °C for 18 h, and the CFU in respective lanes were counted manually.

94

95 **Supplementary information 4: Efficiency of NeACT against internalized *S. aureus* in epithelial** 96 **cells**

97 The cytotoxicity of NeACT was tested in Caco-2 cells ¹². Briefly, the Caco-2 cells were
98 cultured in a 96-well plate (2×10^4 cells/well) until confluent and exposed to increasing
99 concentrations of the particles (3.90-250 μ g/mL) suspended in DMEM media. After 24 h of
100 incubation, resazurin of 50 μ g/mL (prepared with DMEM) was added and incubated for 4 h. The
101 fluorescence was measured at 530/590 nm using a plate reader. Cells without treatment and cells
102 treated with silver nitrate (AgNO_3) were considered as negative and positive controls, respectively.

103

104 The efficiency of NeACT to target internalized *S. aureus* in epithelial cells (Caco-2) was
105 determined by a pre-established protocol with modifications ^{8, 13}. Confluent Caco-2 cells (2×10^4
106 cells/well) were exposed to Sa1158c culture maintained at 1.5×10^8 cells/mL in a 96-well plate and
107 incubated for an hour. The cells were washed using PBS (4 °C) and subjected to gentamicin (10
108 μ g/mL) for 30 min. The extracellular gentamicin was removed by washing, followed by incubation
109 with DMEM for 4 h to establish the intracellular infection model. The particles at incremental
110 concentrations (3.90-15.62 μ g/mL) were added, and the plates were incubated for 24 h in a cell
111 culture humidified incubator at 37 °C, with 5% CO_2 . Subsequently, the cells were washed using
112 PBS and lysed using 0.5% (v/v) of Triton X-100. CFU of viable intracellular bacteria were
113 enumerated using the drop culture method as detailed earlier in section 5.2.8. Sa25923 was used as
114 a reference strain, and cells infected with bacteria without treatment were considered as controls.

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